

Comments /
Suggestions

1	686.4	55.3	886	1	US-08-469-427A-1	Sequence 1, Appli
2	686.4	55.3	886	2	US-08-609-443B-1	Sequence 1, Appli
3	686.4	55.3	886	2	US-08-569-063C-1	Sequence 1, Appli
4	686.4	55.3	886	3	US-08-851-896-1	Sequence 1, Appli
5	609.6	49.1	624	2	US-08-609-443B-12	Sequence 12, Appl
6	609.6	49.1	624	2	US-08-569-063C-12	Sequence 12, Appl
7	609.6	49.1	624	3	US-08-851-896-12	Sequence 12, Appl
8	547.6	44.1	1099	4	US-09-949-016-1545	Sequence 1545, Ap
9	515.2	41.5	624	2	US-08-609-443B-14	Sequence 14, Appl
10	515.2	41.5	624	2	US-08-569-063C-14	Sequence 14, Appl
11	515.2	41.5	624	3	US-08-851-896-14	Sequence 14, Appl
12	441.2	35.5	565	1	US-08-469-427A-4	Sequence 4, Appli
13	441.2	35.5	565	2	US-08-609-443B-4	Sequence 4, Appli
14	441.2	35.5	565	2	US-08-569-063C-4	Sequence 4, Appli
15	441.2	35.5	565	3	US-08-851-896-4	Sequence 4, Appli
16	412.6	33.2	591	1	US-08-469-427A-6	Sequence 6, Appli
17	412.6	33.2	591	2	US-08-609-443B-6	Sequence 6, Appli
18	412.6	33.2	591	2	US-08-569-063C-6	Sequence 6, Appli
19	412.6	33.2	591	3	US-08-851-896-6	Sequence 6, Appli
20	382.2	30.8	405	1	US-08-469-427A-8	Sequence 8, Appli
21	382.2	30.8	405	2	US-08-609-443B-8	Sequence 8, Appli
22	382.2	30.8	405	2	US-08-569-063C-8	Sequence 8, Appli
23	382.2	30.8	405	3	US-08-851-896-8	Sequence 8, Appli
24	355.6	28.6	570	1	US-08-469-427A-10	Sequence 10, Appl
25	355.6	28.6	570	2	US-08-609-443B-10	Sequence 10, Appl
26	355.6	28.6	570	2	US-08-569-063C-10	Sequence 10, Appl
27	355.6	28.6	570	3	US-08-851-896-10	Sequence 10, Appl
28	207.4	16.7	7386	4	US-09-949-016-13287	Sequence 13287, A
29	92.2	7.4	3583	4	US-09-976-594-921	Sequence 921, App
c 30	92.2	7.4	5163	4	US-09-919-039-166	Sequence 166, App
31	91.6	7.4	426	4	US-09-884-050-1	Sequence 1, Appli
32	91.6	7.4	444	3	US-09-392-932-6	Sequence 6, Appli
33	91.6	7.4	444	4	US-09-574-708A-1	Sequence 1, Appli
34	91.6	7.4	444	4	US-09-392-931-1	Sequence 1, Appli
35	91.6	7.4	456	5	PCT-US95-10973A-88	Sequence 88, Appl
36	91.6	7.4	467	5	PCT-US95-10973A-86	Sequence 86, Appl
37	91.6	7.4	473	3	US-08-718-904-1	Sequence 1, Appli
38	91.6	7.4	473	4	US-09-449-249-1	Sequence 1, Appli
39	91.6	7.4	473	5	PCT-US95-10973A-25	Sequence 25, Appl
40	91.6	7.4	495	4	US-09-244-583-25	Sequence 25, Appl
41	91.6	7.4	495	4	US-09-037-983C-14	Sequence 14, Appl
42	91.6	7.4	498	6	5194596-20	Patent No. 5194596
43	91.6	7.4	498	6	5194596-20	Patent No. 5194596
44	91.6	7.4	516	3	US-08-784-551C-1	Sequence 1, Appli
45	91.6	7.4	516	3	US-09-392-932-7	Sequence 7, Appli

ALIGNMENTS

RESULT 1

US-08-469-427A-1

; Sequence 1, Application US/08469427A

; Patent No. 5607918

; GENERAL INFORMATION:

; APPLICANT: Eriksson, Ulf

; APPLICANT: Olofsson, Birgitta

; APPLICANT: Alitalo, Kari

; APPLICANT: Pajusola, Katri

; TITLE OF INVENTION: VASCULAR ENDOTHELIAL GROWTH FACTOR-B AND

; TITLE OF INVENTION: DNA CODING THEREFOR

SCORE Search Results Details for Application 087 and Search Result us-08-765-588a-16.rnpb

[Score Home Page](#) [Retrieve Application List](#) [SCORE System Overview](#) [SCORE FAQ](#) [Comments / Sugg](#)

This page gives you Search Results detail for the Application 08765588 and Search Result us-08-765-588a-16.rnpb.

[start](#)

[Go Back to](#)

GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM nucleic - nucleic search, using sw model

Run on: March 11, 2005, 04:17:05 ; Search time 724 Seconds
(without alignments)
10204.516 Million cell updates/sec

Title: US-08-765-588A-16
Perfect score: 1242
Sequence: 1 gcacgagctcaggccgctcgc.....aaaaaaaaaaaaaaaaaaaaa 1242

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 5537552 seqs, 2974263231 residues

Total number of hits satisfying chosen parameters: 11075104

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_NA:*

- 1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
- 2: /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
- 3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
- 4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
- 5: /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
- 6: /cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
- 7: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
- 8: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
- 9: /cgn2_6/ptodata/1/pubpna/US09A_PUBCOMB.seq:*
- 10: /cgn2_6/ptodata/1/pubpna/US09B_PUBCOMB.seq:*
- 11: /cgn2_6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*
- 12: /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
- 13: /cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
- 14: /cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
- 15: /cgn2_6/ptodata/1/pubpna/US10C_PUBCOMB.seq:*
- 16: /cgn2_6/ptodata/1/pubpna/US10D_PUBCOMB.seq:*
- 17: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*
- 18: /cgn2_6/ptodata/1/pubpna/US10F_PUBCOMB.seq:*
- 19: /cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*

20: /cgn2_6/ptodata/1/pubpna/US11_NEW_PUB.seq:*
 21: /cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
 22: /cgn2_6/ptodata/1/pubpna/US60_PUBCOMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	% Query		Length	DB	ID	Description
	Score	Match				
1	1242	100.0	1242	9	US-09-349-954A-16	Sequence 16, Appl
2	1242	100.0	1242	9	US-09-907-007-16	Sequence 16, Appl
3	1242	100.0	1242	17	US-10-673-708-16	Sequence 16, Appl
4	686.4	55.3	886	10	US-09-961-756-1	Sequence 1, Appli
5	623.4	50.2	1172	19	US-10-278-698-68	Sequence 68, Appl
6	623.4	50.2	1172	19	US-10-278-698-582	Sequence 582, App
7	623.4	50.2	1181	16	US-10-136-819-10	Sequence 10, Appl
8	609.6	49.1	624	10	US-09-961-756-12	Sequence 12, Appl
9	595.2	47.9	1094	9	US-09-349-954A-3	Sequence 3, Appli
10	595.2	47.9	1094	9	US-09-907-007-3	Sequence 3, Appli
11	595.2	47.9	1094	17	US-10-673-708-3	Sequence 3, Appli
12	595.2	47.9	1094	18	US-10-220-324A-1	Sequence 1, Appli
13	571.8	46.0	755	15	US-10-262-538-21	Sequence 21, Appl
14	571.8	46.0	755	15	US-10-007-926A-466	Sequence 466, App
15	571.8	46.0	755	17	US-10-174-128-4	Sequence 4, Appli
16	571.8	46.0	755	18	US-10-669-176-21	Sequence 21, Appl
17	515.2	41.5	624	9	US-09-912-436-3	Sequence 3, Appli
18	515.2	41.5	624	10	US-09-961-756-14	Sequence 14, Appl
19	515.2	41.5	624	18	US-10-384-339C-122	Sequence 122, App
20	515.2	41.5	624	18	US-10-772-927A-3	Sequence 3, Appli
21	513.4	41.3	5695	9	US-09-912-436-10	Sequence 10, Appl
22	512.6	41.3	663	9	US-09-244-694-19	Sequence 19, Appl
23	507	40.8	5695	9	US-09-912-436-9	Sequence 9, Appli
24	503.6	40.5	666	9	US-09-244-694-1	Sequence 1, Appli
25	456.8	36.8	491	9	US-09-244-694-140	Sequence 140, App
26	441.2	35.5	565	10	US-09-961-756-4	Sequence 4, Appli
27	432.8	34.8	910	9	US-09-349-954A-9	Sequence 9, Appli
28	432.8	34.8	910	9	US-09-907-007-9	Sequence 9, Appli
29	432.8	34.8	910	17	US-10-673-708-9	Sequence 9, Appli
30	412.6	33.2	591	10	US-09-961-756-6	Sequence 6, Appli
31	404.4	32.6	1154	17	US-10-264-049-846	Sequence 846, App
32	397	32.0	412	9	US-09-244-694-83	Sequence 83, Appl
33	397	32.0	415	9	US-09-244-694-84	Sequence 84, Appl
34	392.8	31.6	993	9	US-09-349-954A-5	Sequence 5, Appli
35	392.8	31.6	993	9	US-09-907-007-5	Sequence 5, Appli
36	392.8	31.6	993	17	US-10-673-708-5	Sequence 5, Appli
37	392.8	31.6	993	18	US-10-220-324A-3	Sequence 3, Appli
38	382.2	30.8	405	10	US-09-961-756-8	Sequence 8, Appli
39	375.8	30.3	410	9	US-09-244-694-141	Sequence 141, App
40	370	29.8	381	9	US-09-244-694-36	Sequence 36, Appl
41	355.6	28.6	570	9	US-09-795-006A-116	Sequence 116, App
42	355.6	28.6	570	10	US-09-961-756-10	Sequence 10, Appl
43	355.6	28.6	570	18	US-10-772-927A-1	Sequence 1, Appli
44	355.2	28.6	567	9	US-09-912-436-1	Sequence 1, Appli
45	353.2	28.4	5614	9	US-09-912-436-7	Sequence 7, Appli

ALIGNMENTS

SCORE Search Results Details for Application 08765588 and Search Result us-08-765-588a-16.rnpm.

[Score Home](#)
[Page](#)

Retrieve Application List

SCORE System Overview

SCORE
FAO

Comments /
Suggestions

This page gives you Search Results detail for the Application 08765588 and Search Result us-08-765-588a-16.rnrm.

start

[Go Back to previous page](#)

GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM nucleic - nucleic search, using sw model

```
Run on:      March 11, 2005, 03:12:43 ; Search time 4480 Seconds
              (without alignments)
              11317.460 Million cell updates/sec
```

```
Title:          US-08-765-588A-16
Perfect score: 1242
Sequence:      1 qcacgaqctcagggcctcg.....aaaaaaaaaaaaaaaaaaaaa 1242
```

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 45554873 seqs, 20411521753 residues

Total number of hits satisfying chosen parameters: 91109746

Minimum DB seq length: 0

```
Maximum DB seq length: 2000000000
```

```
Post-processing: Minimum Match 0%
                  Maximum Match 100%
                  Listing first 45 summaries
```

```
Database : Pending_Patents_NA_Main:*
1: /cgn2_6/ptodata/1/pna/PCTUS1_COMB.seq:*
2: /cgn2_6/ptodata/1/pna/PCTUS2_COMB.seq:*
3: /cgn2_6/ptodata/1/pna/PCTUS_COMB.seq:*
4: /cgn2_6/ptodata/1/pna/US06_COMB.seq:*
5: /cgn2_6/ptodata/1/pna/US07_COMB.seq:*
6: /cgn2_6/ptodata/1/pna/US080_COMB.seq:*
7: /cgn2_6/ptodata/1/pna/US081_COMB.seq:*
8: /cgn2_6/ptodata/1/pna/US082_COMB.seq:*
9: /cgn2_6/ptodata/1/pna/US083_COMB.seq:*
10: /cgn2_6/ptodata/1/pna/US084_COMB.seq:*
11: /cgn2_6/ptodata/1/pna/US085_COMB.seq:*
12: /cgn2_6/ptodata/1/pna/US086_COMB.seq:*
13: /cgn2_6/ptodata/1/pna/US087_COMB.seq:*
14: /cgn2_6/ptodata/1/pna/US088_COMB.seq:*
15: /cgn2_6/ptodata/1/pna/US089_COMB.seq:*
16: /cgn2_6/ptodata/1/pna/US090_COMB.seq:*
```

17: /cgn2_6/ptodata/1/pna/US091_COMB.seq:*
18: /cgn2_6/ptodata/1/pna/US092A_COMB.seq:*
19: /cgn2_6/ptodata/1/pna/US092B_COMB.seq:*
20: /cgn2_6/ptodata/1/pna/US093A_COMB.seq:*
21: /cgn2_6/ptodata/1/pna/US093B_COMB.seq:*
22: /cgn2_6/ptodata/1/pna/US094_COMB.seq:*
23: /cgn2_6/ptodata/1/pna/US095A_COMB.seq:*
24: /cgn2_6/ptodata/1/pna/US095B_COMB.seq:*
25: /cgn2_6/ptodata/1/pna/US095C_COMB.seq:*
26: /cgn2_6/ptodata/1/pna/US095D_COMB.seq:*
27: /cgn2_6/ptodata/1/pna/US096A_COMB.seq:*
28: /cgn2_6/ptodata/1/pna/US096B_COMB.seq:*
29: /cgn2_6/ptodata/1/pna/US096C_COMB.seq:*
30: /cgn2_6/ptodata/1/pna/US096D_COMB.seq:*
31: /cgn2_6/ptodata/1/pna/US096E_COMB.seq:*
32: /cgn2_6/ptodata/1/pna/US097A_COMB.seq:*
33: /cgn2_6/ptodata/1/pna/US097B_COMB.seq:*
34: /cgn2_6/ptodata/1/pna/US097C_COMB.seq:*
35: /cgn2_6/ptodata/1/pna/US098A_COMB.seq:*
36: /cgn2_6/ptodata/1/pna/US098B_COMB.seq:*
37: /cgn2_6/ptodata/1/pna/US098C_COMB.seq:*
38: /cgn2_6/ptodata/1/pna/US098D_COMB.seq:*
39: /cgn2_6/ptodata/1/pna/US099A_COMB.seq:*
40: /cgn2_6/ptodata/1/pna/US099B_COMB.seq:*
41: /cgn2_6/ptodata/1/pna/US099C_COMB.seq:*
42: /cgn2_6/ptodata/1/pna/US099D_COMB.seq:*
43: /cgn2_6/ptodata/1/pna/US099E_COMB.seq:*
44: /cgn2_6/ptodata/1/pna/US099F_COMB.seq:*
45: /cgn2_6/ptodata/1/pna/US099G_COMB.seq:*
46: /cgn2_6/ptodata/1/pna/US100A_COMB.seq:*
47: /cgn2_6/ptodata/1/pna/US100B_COMB.seq:*
48: /cgn2_6/ptodata/1/pna/US101A_COMB.seq:*
49: /cgn2_6/ptodata/1/pna/US101B_COMB.seq:*
50: /cgn2_6/ptodata/1/pna/US102A_COMB.seq:*
51: /cgn2_6/ptodata/1/pna/US102B_COMB.seq:*
52: /cgn2_6/ptodata/1/pna/US103A_COMB.seq:*
53: /cgn2_6/ptodata/1/pna/US103B_COMB.seq:*
54: /cgn2_6/ptodata/1/pna/US104A_COMB.seq:*
55: /cgn2_6/ptodata/1/pna/US104B_COMB.seq:*
56: /cgn2_6/ptodata/1/pna/US105A_COMB.seq:*
57: /cgn2_6/ptodata/1/pna/US105B_COMB.seq:*
58: /cgn2_6/ptodata/1/pna/US106A_COMB.seq:*
59: /cgn2_6/ptodata/1/pna/US107A_COMB.seq:*
60: /cgn2_6/ptodata/1/pna/US107B_COMB.seq:*
61: /cgn2_6/ptodata/1/pna/US107C_COMB.seq:*
62: /cgn2_6/ptodata/1/pna/US107D_COMB.seq:*
63: /cgn2_6/ptodata/1/pna/US108A_COMB.seq:*
64: /cgn2_6/ptodata/1/pna/US108B_COMB.seq:*
65: /cgn2_6/ptodata/1/pna/US109A_COMB.seq:*
66: /cgn2_6/ptodata/1/pna/US109B_COMB.seq:*
67: /cgn2_6/ptodata/1/pna/US109C_COMB.seq:*
68: /cgn2_6/ptodata/1/pna/US110_COMB.seq:*
69: /cgn2_6/ptodata/1/pna/US6000_COMB.seq:*
70: /cgn2_6/ptodata/1/pna/US6001_COMB.seq:*
71: /cgn2_6/ptodata/1/pna/US6002_COMB.seq:*
72: /cgn2_6/ptodata/1/pna/US6003_COMB.seq:*
73: /cgn2_6/ptodata/1/pna/US6004_COMB.seq:*
74: /cgn2_6/ptodata/1/pna/US6005_COMB.seq:*
75: /cgn2_6/ptodata/1/pna/US6006_COMB.seq:*
76: /cgn2_6/ptodata/1/pna/US6007_COMB.seq:*
77: /cgn2_6/ptodata/1/pna/US6008_COMB.seq:*

SCORE Search Results Details for Application 08765588 and Search Result us-08-765-588a-16.rnpn.

Score Home	Retrieve Application	SCORE System	SCORE	Comments /
Page	List	Overview	FAQ	Suggestions

This page gives you Search Results detail for the Application 08765588 and Search Result us-08-765-588a-16.rnpn.

[start](#)

[Go Back to previous page](#)

GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM nucleic - nucleic search, using sw model

Run on: March 11, 2005, 04:13:05 ; Search time 1462 Seconds
(without alignments)
624.206 Million cell updates/sec

Title: US-08-765-588A-16
Perfect score: 1242
Sequence: 1 gcacgagctcaggccgtcgc.....aaaaaaaaaaaaaaaaaaaaa 1242

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 5522993 seqs, 367386938 residues

Total number of hits satisfying chosen parameters: 11045986

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Pending_Patents_NA_New:*

- 1: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*
- 2: /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*
- 3: /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*
- 4: /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
- 5: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
- 6: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*
- 7: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq2:*
- 8: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq:*
- 9: /cgn2_6/ptodata/1/pna/US60_NEW_COMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

%

Day : Thursday
Date: 6/15/2006

Time: 10:02:02

 **PALM INTRANET**

Inventor Information for 08/765588

Inventor Name	City	State/Country
HAYWARD, NICHOLAS KIM	PADDINGTON	AUSTRALIA
WEBER, GUNTHER	STOCKHOLM	SWEDEN
GRIMMOND, SEAN	TARANGA	AUSTRALIA
NORDENSKJOLD, MAGNUS	STOCKHOLM	SWEDEN
LARSSON, CATHARINA	STOCKHOLM	SWEDEN

[Appln Info](#)[Contents](#)[Petition Info](#)[Atty/Agent Info](#)[Continuity Data](#)[Foreign Data](#)Search Another: Application# or Patent# PCT / / or PG PUBS # Attorney Docket # Bar Code #

To go back use Back button on your browser toolbar.

Back to [PALM](#) | [ASSIGNMENT](#) | [OASIS](#) | [Home page](#)

Day : Thursday
Date: 6/15/2006

Time: 10:03:13

 **PALM INTRANET**

Inventor Information for 08/569063

Inventor Name	City	State/Country
ERIKSSON, ULF	BALSTA	SWEDEN
OLOFSSON, BIRGITTA	SUNDBYBERG	SWEDEN
ALITALO, KARI	HELSINKI	FINLAND
PAJUSOLA, KATRI	HELSINKI	FINLAND

[Appln Info](#)[Contents](#)[Petition Info](#)[Atty/Agent Info](#)[Continuity Data](#)[Foreign Data](#)Search Another: Application# [Search](#) or Patent# [Search](#)PCT / / [Search](#) or PG PUBS # [Search](#)Attorney Docket # [Search](#)Bar Code # [Search](#)

To go back use Back button on your browser toolbar.

Back to [PALM](#) | [ASSIGNMENT](#) | [OASIS](#) | [Home page](#)